

AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions, and listing, of claims in the application:

Listing of Claims

1. (currently amended) An isolated nucleic acid sequence which encodes a polypeptide with desaturase activity, selected from the following group:
 - a) a nucleic acid sequence with the sequence shown in SEQ ID NO: 1,
 - b) a nucleic acid sequence which, as a result of the degeneracy of the genetic code, is are derived from the nucleic acid sequence shown in SEQ ID NO: 1,
 - c) derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequence shown in SEQ ID NO: 2 and which have at least 90% homology at amino acid level without reducing the enzymatic activity of the polypeptides to less than 10% of the activity of the polypeptides with the amino acid sequence shown in SEQ ID NO: 2.
- 2-3 (canceled)
- 4 (previously presented). A nucleic acid construct comprising the nucleic acid sequence as claimed in claim 1, where the nucleic acid sequence is linked to one or more regulatory sequences.
- 5 (previously presented). A vector comprising the nucleic acid sequence as claimed in claim 1, or a nucleic acid construct comprising said nucleic acid sequence linked to one or more regulatory sequences.
- 6 (previously presented). A non-human organism comprising the nucleic acid sequence as claimed in claim 1 or at least one nucleic acid construct comprising said nucleic acid sequence linked to one or more regulatory sequences.

- 7 (previously presented). The non-human organism as claimed in claim 6, which is a plant, a microorganism or an animal.
- 8 (currently amended). A transgenic plant comprising the nucleic acid sequence as claimed in claim 1, wherein said nucleic acid sequence is a nucleic acid construct comprising said nucleic acid sequence linked to one or more regulatory sequences.
- 9 (previously presented). A process for the preparation of saturated or unsaturated fatty acids, which comprises introducing the nucleic acid sequence as claimed in claim 1 or a nucleic acid construct comprising said nucleic acid linked to one or more regulatory sequences into an organism, growing this organism, isolating oil contained in the organism and liberating fatty acids contained in the oil.
- 10 (previously presented). A process for the preparation of triglycerides with an increased content of unsaturated fatty acids, which comprises introducing the nucleic acid sequence as claimed in claim 1 or a nucleic acid construct comprising said nucleic acid linked to one or more regulatory sequences into an organism, growing this organism and isolating the oil contained in the organism.
- 11 (canceled)
- 12 (currently amended). A process for the preparation of triglycerides with an increased content of saturated fatty acids, which comprises introducing the nucleic acid sequence as claimed in claim 1, wherein said nucleic acid sequence is a nucleic acid construct comprising said nucleic acid sequence linked to one or more regulatory sequences into an organism, growing this organism and isolating oil contained in the organism.
- 13 (previously presented). The process as claimed in claim 9, wherein the unsaturated fatty acid has an increased calendulic acid content.

14 (previously presented). The process claimed in claim 9, wherein the organism is one of a plant and a microorganism.

15-23 (canceled)

24 (previously presented). The isolated nucleic acid sequence which encodes a polypeptide with desaturase activity of claim 1, wherein said derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequence shown in SEQ ID NO: 2 and which have at least 90% homology at amino acid level without reducing the desaturase activity of the polypeptides to less than 20% of the desaturase activity of the polypeptides with the amino acid sequence shown in SEQ ID NO: 2.

25 (previously presented). The isolated nucleic acid sequence which encodes a polypeptide with desaturase activity of claim 1, wherein said derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequence shown in SEQ ID NO: 2 and which have at least 90% homology at amino acid level without reducing the desaturase activity of the polypeptides to less than 30% of the desaturase activity of the polypeptides with the amino acid sequence shown in SEQ ID NO: 2.

26 (previously presented). The isolated nucleic acid sequence which encodes a polypeptide with desaturase activity of claim 1, wherein said derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequence shown in SEQ ID NO: 2 and which have at least 95% homology at amino acid level without reducing the desaturase activity of the polypeptides to less than 20% of the desaturase activity of the polypeptides with the amino acid sequence shown in SEQ ID NO: 2.

27 (currently amended). The ~~desaturase of claim 1~~ isolated nucleic acid sequence which encodes the polypeptide with desaturase activity of claim 1 wherein said ~~enzyme polypeptide~~ causes a regiospecific shift of a *cis* double bond in position C9 to a *trans* double bond in position C10 and introduces a *trans* double bond at position C8.

28 (previously presented). An isolated nucleic acid comprising SEQ ID NO: 1.

29-30 (canceled)